

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/582,918
Source: FWP
Date Processed by STIC: 6/22/06

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

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Revised 01/10/06



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/582,918

DATE: 06/22/2006

TIME: 13:10:32

Input Set : E:\seqlist.txt

Output Set: N:\CRF4\06222006\J582918.raw

3 <110> APPLICANT: BASF Aktiengesellschaft
 5 <120> TITLE OF INVENTION: P EF-TU-Expressionseinheiten
 7 <130> FILE REFERENCE: PF 55183/Mec
 W--> 8 <140> CURRENT APPLICATION NUMBER: 20030320
 C--> 10 <141> CURRENT FILING DATE: 2006-06-14
 12 <160> NUMBER OF SEQ ID NOS: 42

use English in a
 U.S.

this
 is NOT the
 current application number

ERRORED SEQUENCES

2271 <210> SEQ ID NO: 42
 2272 <211> LENGTH: 7
 2273 <212> TYPE: DNA
 2274 <213> ORGANISM: Corynebacterium glutamicum
 2276 <220> FEATURE:
 2277 <223> OTHER INFORMATION: RIBOSOMALE
 2279 <400> SEQUENCE: 42
 2280 aggagga
 E--> 2283 PF 55183
 E--> 2285 5
 E--> 2289 PF 55183

Does Not Comply
 Corrected Diskette Needed

use English

delete at end of file

Please
 use English
 throughout the
 submitted file

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/582,918

DATE: 06/22/2006

TIME: 13:10:34

Input Set : E:\seqlist.txt

Output Set: N:\CRF4\06222006\J582918.raw

L:8 M:283 W: Missing Blank Line separator, <140> field identifier
L:8 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2283 M:254 E: No. of Bases conflict, LENGTH:Input:55183 Counted:9 SEQ:42
L:2283 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=42
L:2289 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:2289 M:252 E: No. of Seq. differs, <211> LENGTH:Input:7 Found:11 SEQ:42